

SEQUENCE LISTING

<110> Smith, Theresa H.
 <120> PRO-INFLAMMATORY FIBRINOPEPTIDE
 <130> US 1257/01 (VA)
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 <223> STRANDEDNESS: single
 TOPOLOGY: both

<400> SEQ ID NO: 1
 Ala Asp Ser Gly Glu Gly Asp Phe Leu Ala Glu Gly Gly Gly Val
 5 10 15
 Arg Gly Pro Arg Val Val Glu Arg His Gln Ser Ala Cys Lys Asp
 20 25 30
 Ser Asp Trp Pro Phe Cys Ser Asp Glu Asp Trp Asn Tyr Lys Cys
 35 40 45
 Pro Ser Gly Cys Arg Met Lys Gly Leu Ile Asp Glu Val Asn Gln
 50 55 60
 Asp Phe Thr Asn Arg Ile Asn Lys Leu Lys Asn Ser Leu Phe Glu
 65 70 75
 Tyr Gln Lys Asn Asn Lys Asp Ser His Ser Leu Thr Thr Asn Ile
 80 85 90
 Met Glu Ile Leu Arg Gly Asp Phe Ser Ser Ala Asn Asn Arg Asp
 95 100 105
 Asn Thr Tyr Asn Arg Val Ser Glu Asp Leu Arg Ser Arg Ile Glu
 110 115 120
 Val Leu Lys Arg Lys Val Ile Glu Lys Val Gln His Ile Gln Leu
 125 130 135
 Leu Gln Lys Asn Val Arg Ala Gln Leu Val Asp Met Lys Arg Leu
 140 145 150
 Glu Val Asp Ile Asp Ile Lys Ile Arg Ser Cys Arg Gly Ser Cys
 155 160 165
 Ser Arg Ala Leu Ala Arg Glu Val Asp Leu Lys Asp Tyr Glu Asp
 170 175 180
 Gln Gln Lys Gln Leu Glu Gln Val Ile Ala Lys Asp Leu Leu Pro
 185 190 195

Ser Arg Asp Arg	Gln His Leu Pro Leu	Ile Lys Met Lys Pro	Val
200	205		210
Pro Asp Leu Val	Pro Gly Asn Phe Lys	Ser Gln Leu Gln Lys	Val
215	220		225
Pro Pro Glu Trp	Lys Ala Leu Thr Asp	Met Pro Gln Met Arg	Met
230	235		240
Glu Leu Glu Arg	Pro Gly Gly Asn Glu	Ile Thr Arg Gly Gly	Ser
245	250		255
Thr Ser Tyr Gly	Thr Gly Ser Glu Thr	Glu Ser Pro Arg Asn	Pro
260	265		270
Ser Ser Ala Gly	Ser Trp Asn Ser Gly	Ser Ser Gly Pro Gly	Ser
275	280		285
Thr Gly Asn Arg	Asn Pro Gly Ser Ser	Gly Thr Gly Gly Thr	Ala
290	295		300
Thr Trp Lys Pro	Gly Ser Ser Gly Pro	Gly Ser Thr Gly Ser	Trp
305	310		315
Asn Ser Gly Ser	Ser Gly Thr Gly Ser	Thr Gly Asn Gln Asn	Pro
320	325		330
Gly Ser Pro Arg	Pro Gly Ser Thr Gly	Thr Trp Asn Pro Gly	Ser
335	340		345
Ser Glu Arg Gly	Ser Ala Gly His Trp	Thr Ser Glu Ser Ser	Val
350	355		360
Ser Gly Ser Thr	Gly Gln Trp His Ser	Glu Ser Gly Ser Phe	Arg
365	370		375
Pro Asp Ser Pro	Gly Ser Gly Asn Ala	Arg Pro Asn Asn Pro	Asp
380	385		390
Trp Gly Thr Phe	Glu Glu Val Ser Gly	Asn Val Ser Pro Gly	Thr
395	400		405
Arg Arg Glu Tyr	His Thr Glu Lys Leu	Val Thr Ser Lys Gly	Asp
410	415		420
Lys Glu Leu Arg	Thr Gly Lys Glu Lys	Val Thr Ser Gly Ser	Thr
425	430		435
Thr Thr Thr Arg	Arg Ser Cys Ser Lys	Thr Val Thr Lys Thr	Val
440	445		450
Ile Gly Pro Asp	Gly His Lys Glu Val	Thr Lys Glu Val Val	Thr
455	460		465
Ser Glu Asp Gly	Ser Asp Cys Pro Glu	Ala Met Asp Leu Gly	Thr
470	475		480
Leu Ser Gly Ile	Gly Thr Leu Asp Gly	Phe Arg His Arg His	Pro
485	490		495

